

Remarks:

Reconsideration of the application is requested.

Please amend the specification as directed above. Included is a Sequence Listing to be added to the end of the specification. Support for the Sequence Listing can be found on page 8 of the originally filed specification.

I hereby state that the submission, filed in accordance with 37 CFR 1.821(g), herein does not include new matter.

A copy of the Notice is attached.

In view of the foregoing, examination of the application is solicited.

Petition for extension is herewith made. The extension fee for response within a period of one month pursuant to Section 1.136(a) in the amount of \$55 in accordance with Section 1.17 is enclosed herewith.

Please charge any other fees that might be due with respect to Sections 1.16 and 1.17 to the Deposit Account of Lerner and Greenberg, P.A., No. 12-1099.

Respectfully submitted,

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For Applicant

LDP:cgm

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Version with Markings to Show Changes Made:In the Specification:

Replace the paragraph beginning at page 7, line 14, with the following:

-- The following is an example of how this procedure may be used to detect an HIV viral load. Intravenously drawn blood is collected and 1 milliliter is diluted into a solution of RNA preservative solution. The RNA is purified out of the solution by the usual biochemical protocols with the final step eluting from a silica spin column. The RNA is eluted with RNA free water, 40 micro liters final volume, and placed into a PCR reaction tube. To this solution will be added 20 units AMV Reverse Transcriptase, 10X buffer [suitable for use with the particular PCR enzymes], and 3 units of Hot Start DNA Polymerase, 25 micromoles each of dUTP, dATP, dCTP, & dGTP. The primers are labeled as follows and are constructed in a 5' to 3' direction suitable for RT-PCR of HIV. SEQ ID NO:1 HIV GAG Forward FAM [5']-ATAATCCACCTAATCCCAGTAGGAGAAAT- [3'], mp= 78C, and SEQ ID NO:2 GAG Reverse BIOTIN [5']-TTTGGTCCTTGTCTTATGTCCAGAATG with a mp= 76C and were added to the reaction at a level of one micro molar. All of the reagents that are added to the RNA can be stored as a concentrate solution so as to impart an automated ability to the assay. An arbitrary 10 micro liters of reagent could be added to the 40 micro liters of RNA to make a 50 micro liter

RT-PCR reaction. The reaction conditions are as follows: the Reverse Transcription step is allowed to occur over the course of one hour with a starting temperature of 45C and reaching a final temperature of 65C at an increase of 1 C per 3 minute, the "Hot Start" is allowed to proceed at 95 C for 15 minutes to liberate the DNA Polymerase activity, and then the following temperatures will be maintained for 40 cycles [1] 94 C for 30 seconds, [2] 55 C for 30 seconds, & [3] 72 C for 30 seconds. After the PCR, the reactants are pipetted into a strepavidin-coated plate suitable for use with a 96 well fluorometer. The binding reaction of Biotin to strepavidin is allowed to occur for a minimum of 30 minutes. The well is rinsed 3 times with Tween Tris Buffered Saline for three times [this step can be done manually or automatically]. The plate is then read for the amount of fluorescence and this should be directly proportional to the amount of virus isolated. In this particular reaction, FAM will be a 5' labeled Fluorescein for the detection of fluorescence and the Biotin, 18-space linker, will be used for capturing the Amplicons onto a strepavidin coated plate. It does not matter whether the forward or reverse primer gets the capture or detection agent as the Amplicons will still have the ability to be read and their construction is not critical. The ability of the Amplicons to be simultaneously captured, washed, and subsequently detected is what is novel about the modifications of this reaction and analysis mechanism. Although this

procedure does NOT lend itself to a real-time assay, it does NOT require the expensive apparatus either. The design of this assay lends itself to automation for high throughput screening at a relatively low cost with reagents readily available.--

In addition, add the attached sequence listing after the last page of the specification.